p#8

1653

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/664,326

DATE: 04/04/2002
TIME: 15:16:31

Input Set : A:\24811693.app
Output Set: N:\CRF3\04042002\I664326.raw


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      3 <110> APPLICANT: HABERMANN, PAUL
      4       BENDER, RUDOLF
      6 <120> TITLE OF INVENTION: SIGNAL SEQUENCES FOR PREPARING LEU-HIRUDIN BY SECRETION
      7       BY E. COLI INTO THE CULTURE MEDIUM
      9 <130> FILE REFERENCE: 02481.1693
C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/664,326
C--> 12 <141> CURRENT FILING DATE: 2000-09-18
     14 <160> NUMBER OF SEQ ID NOS: 33
     16 <170> SOFTWARE: PatentIn Ver. 2.1
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     45 <213> ORGANISM: Artificial Sequence
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     57 <213> ORGANISM: Artificial Sequence
     59 <220> FEATURE:
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     67 <211> LENGTH: 56
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71 <220> FEATURE:
72 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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88 a 61
91 <210> SEQ ID NO: 7
92 <211> LENGTH: 59
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94 <213> ORGANISM: Artificial Sequence
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108 <220> FEATURE:
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113 gca 63
116 <210> SEQ ID NO: 9
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119 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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136 <400> SEQUENCE: 10
137 ctgatcccgt tctttgcagc gttctgcctg ccggttttcg cgcttacgta tactgactgc 60
138 a 61

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<220> FEATURE:
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tgca 64

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224 atggc 65
227 <210> SEQ ID NO: 18
228 <211> LENGTH: 55
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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236 tggtttcagc tttagtaagc ggggttgcat ttgctcttac gtatactgac tgcac 55
239 <210> SEQ ID NO: 19
240 <211> LENGTH: 60
241 <212> TYPE: DNA
242 <213> ORGANISM: Artificial Sequence
244 <220> FEATURE:
245 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
247 <400> SEQUENCE: 19
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253 <212> TYPE: DNA
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256 <220> FEATURE:
257 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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264 <211> LENGTH: 57
265 <212> TYPE: DNA
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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275 <210> SEQ ID NO: 22
276 <211> LENGTH: 227
277 <212> TYPE: DNA
278 <213> ORGANISM: Unknown Organism
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Description of Unknown Organism: Hirudin-encoding
282 DNA sequence
284 <220> FEATURE:

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290 Leu Thr Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys
291   1               5                  10                  15
293 gaa gga tct aac gtt tgc ggc cag ggt aac aaa tgc atc ctt gga tcc   96
294 Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser
295              20                  25                  30
297 gac ggt gaa aag aac cag tgc gtt act ggc gaa ggt acc ccg aaa ccg   144
298 Asp Gly Glu Lys Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
299          35                  40                  45
301 cag tct cat aac gac ggc gac ttc gaa gag atc cct gag gaa tac ctt   192
302 Gln Ser His Asn Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu
303      50                  55                  60
305 cag taatagagct cgtcgacctg cagcccaagc tt                          227
306 Gln
307  65
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311 <211> LENGTH: 65
312 <212> TYPE: PRT
313 <213> ORGANISM: Unknown Organism
315 <220> FEATURE:
316 <223> OTHER INFORMATION: Description of Unknown Organism: Hirudin-encoded
317       amino acid sequence
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320 Leu Thr Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys
321   1               5                  10                  15
323 Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser
324              20                  25                  30
326 Asp Gly Glu Lys Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
327          35                  40                  45
329 Gln Ser His Asn Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu
330      50                  55                  60
332 Gln
333  65
337 <210> SEQ ID NO: 24
338 <211> LENGTH: 30
339 <212> TYPE: PRT
340 <213> ORGANISM: Unknown Organism
342 <220> FEATURE:
343 <223> OTHER INFORMATION: Description of Unknown Organism: Control:
344       cgtase-Ala-hirudin
346 <400> SEQUENCE: 24
347 Met Lys Arg Asn Arg Phe Phe Asn Thr Ser Ala Ala Ile Ala Ile Ser
348   1               5                  10                  15
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351              20                  25                  30
354 <210> SEQ ID NO: 25
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L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date